



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/721,405
Source: IFW9
Date Processed by STIC: 12/17/03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 4B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/721,405

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 00701/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/721,405

DATE: 12/17/2003

TIME: 07:58:39

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12162003\J721405.raw

3 <110> APPLICANT: Agouron Pharmaceuticals, Inc./ A Pfizer Company
 5 <120> TITLE OF INVENTION: DUAL ASSAY FOR EVALUATING ACTIVITY AND CYTOTOXICITY OF
 6 COMPOUNDS IN THE SAME POPULATION OF CELLS
 8 <130> FILE REFERENCE: PC25522A
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/721,405
 11 <141> CURRENT FILING DATE: 2003-11-24
 13 <150> PRIOR APPLICATION NUMBER: 60/429,382
 14 <151> PRIOR FILING DATE: 2002-11-25
 16 <160> NUMBER OF SEQ ID NOS: 23
 18 <170> SOFTWARE: PatentIn version 3.1
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 936
 22 <212> TYPE: DNA
 23 <213> ORGANISM: Renilla Luciferase Humanized Codons
 25 <400> SEQUENCE: 1

26	atgacctcca	agggtgtacga	ccccgagcag	cgcaagcgca	tgattaccgg	cccccagtgg	60
28	tgggcccgcg	gcaagcagat	gaacgtgctg	gacagcttca	tcaactacta	cgacagcgag	120
30	aagcacgccg	agaacgccgt	gatcttcctg	cacggcaacg	ccgccagctc	ctacctgtgg	180
32	cgccacgtgg	tgcctcacat	cgagcctgtg	gcccgtgca	tcattccctga	cctgatcggc	240
34	atgggcaaga	gcggcaagag	cggaacggc	agctaccgcc	tgctggacca	ctacaagtac	300
36	ctgaccgcct	ggttcgagct	gctgaacctg	cccaagaaga	tcattcttct	gggccacgac	360
38	tggggcgcc	gcctggcctt	ccactacagc	tacgagcacc	aggacaagat	caaggccatc	420
40	gtgcacgccg	agagcgtggt	ggacgtgatc	gagagctggg	acgagtggcc	tgacatcgag	480
42	gaggacatcg	ccctgatcaa	gagcgaggag	ggcgagaaga	tggtgctgga	gaacaacttc	540
44	ttcgttgaga	ccatgctgcc	tagcaagatc	atgcgcaagc	tgagcctga	ggagttcgcc	600
46	gcctacctgg	agcccttcaa	ggagaagggc	gaggtgcgcc	gccctaccct	gagctggcct	660
48	gcgcgagatc	ctctggtgaa	ggcgggcaag	cctgacgtgg	tgcatatcgt	gcgcaactac	720
50	aacgcctacc	tgcgcgccag	cgacgacctg	cccaagatgt	tcattcgagag	cgaccctggc	780
52	ttcttcagca	acgccatcgt	ggaggcgccc	aagaagttcc	ctaaccacca	gttcgtgaag	840
54	gtgaagggcc	tgcaattcag	ccaggaggac	gcccctgacg	agatgggcaa	gtacatcaag	900
56	agcttcgtgg	agcgcgtgct	gaagaacgag	cagtaa			936

59 <210> SEQ ID NO: 2
 60 <211> LENGTH: 936
 61 <212> TYPE: DNA
 62 <213> ORGANISM: Renilla reniformis
 64 <400> SEQUENCE: 2

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67	tgggccagat	gtaaacaat	gaatgttctt	gattcattta	ttaattatta	tgattcagaa	120
69	aaacatgcag	aaaatgctgt	tattttttta	catggtaacg	cgccctcttc	ttattttatg	180
71	cgacatgttg	tgccacatat	tgagccagta	gcgcggtgta	ttataccaga	ccttatttgt	240
73	atgggcaaat	caggcaaatc	tggtaatggt	tcttataggt	tacttgatca	ttacaaatat	300
75	cttactgcat	ggtttgaact	tcttaattta	ccaaagaaga	tcatttttgt	cggccatgat	360
77	tggggtgctt	gttttggcatt	tcatttatagc	tatgagcatc	aagataagat	caaagcaata	420

Does Not Comply
 Corrected Disclosure Needed
 pp 2-5

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Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12162003\J721405.raw

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79 gttcacgctg aaagtgtagt agatgtgatt gaatcatggg atgaatggcc tgatattgaa 480
81 gaagatattg cgttgatcaa atctgaagaa ggagaaaaaa tggttttgga gaataacttc 540
83 ttcgtggaaa ccatgttgcc atcaaaaatc atgagaaagt tagaaccaga agaatttgca 600
85 gcatactctg aaccattcaa agagaaaggt gaagttcgct gtccaacatt atcatggcct 660
87 cgtgaaatcc cgttagttaa aggtggtaaa cctgacgttg tacaaattgt taggaattat 720
89 aatgcttatac tacgtgcaag tgatgattta ccaaaaatgt ttattgaatc ggaccagga 780
91 ttcttttcca atgctattgt tgaaggtgcc aagaagtttc ctaatactga atttgtcaaa 840
93 gtaaaaggtc ttcatttttc gcaagaagat gcacctgatg aaatgggaaa atatatcaaa 900
95 tcgttcgttg agcgagttct caaaaatgaa caataa 936

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98 <210> SEQ ID NO: 3

99 <211> LENGTH: 75

100 <212> TYPE: DNA

101 <213> ORGANISM: Oligonucleotide Template

103 <400> SEQUENCE: 3

104 atgacctcca aggtgtacga ccccgagcag cgcaagcgca tgattaccgg cccccagtgg 60

106 tgggcccgct gcaag 75

109 <210> SEQ ID NO: 4

110 <211> LENGTH: 38

111 <212> TYPE: DNA

112 <213> ORGANISM: Oligonucleotide Primer

114 <400> SEQUENCE: 4

115 gaatcatcta gaatgacctc caaggtgtac gacccgca 38

118 <210> SEQ ID NO: 5

119 <211> LENGTH: 33

120 <212> TYPE: DNA

121 <213> ORGANISM: Oligonucleotide Primer

123 <400> SEQUENCE: 5

124 gttcatgaat tccttgacgc gggccaccca ctg 33

127 <210> SEQ ID NO: 6

128 <211> LENGTH: 99

129 <212> TYPE: DNA

130 <213> ORGANISM: Oligonucleotide Template

132 <400> SEQUENCE: 6

133 gtgctggaca gtttcatcaa ctactacgac agcgagaagc acgccgagaa cgccgtgac 60

135 ttcttgacgc gcaacgccgc cagctctac ctgtggcgc 99

138 <210> SEQ ID NO: 7

139 <211> LENGTH: 100

140 <212> TYPE: DNA

141 <213> ORGANISM: Oligonucleotide Primer

143 <400> SEQUENCE: 7

144 cgctcttgcc catgccgatc aggtcagga tgatgcagcg ggccacaggc tcgatgtgag 60

146 gcaccacgtg ggcacacagg taggagctgg cggcgttgcc 100

149 <210> SEQ ID NO: 8

150 <211> LENGTH: 65

151 <212> TYPE: DNA

152 <213> ORGANISM: Oligonucleotide Primer

154 <400> SEQUENCE: 8

155 gaatcatcta gatgggcccg ctgcaagcag atgaacgtgc tggacagctt catcaactac 60

157 tacga 65

*invalid response - see item 10
on Enr summary
sheet*

RAW SEQUENCE LISTING

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Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12162003\J721405.raw

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163 <213> ORGANISM: Oligonucleotide Primer
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168 cgatcaggtc                                                                    70
171 <210> SEQ ID NO: 10
172 <211> LENGTH: 98
173 <212> TYPE: DNA
174 <213> ORGANISM: Oligonucleotide Template
176 <400> SEQUENCE: 10
177 cgagctgctg aacctgccca agaagatcat cttcgtgggc cacgactggg ggcctgcct      60
179 ggccttcac tacagctacg agcaccagga caagatca                                98
182 <210> SEQ ID NO: 11
183 <211> LENGTH: 99
184 <212> TYPE: DNA
185 <213> ORGANISM: Oligonucleotide template
187 <400> SEQUENCE: 11
188 cctcctcgat gtcaggccac tcgtcccagc tctcgatcac gtccaccacg ctctcggcgt      60
190 gcacgatggc cttgatcttg tcttggtgct cgtagctgt                            99
193 <210> SEQ ID NO: 12
194 <211> LENGTH: 70
195 <212> TYPE: DNA
196 <213> ORGANISM: Oligonucleotide Primer
198 <400> SEQUENCE: 12
199 accgcctgct ggaccactac aagtaoctga ccgcctgggt cgagctgctg aacctgccca      60
201 agaagatcat                                                                    70
204 <210> SEQ ID NO: 13
205 <211> LENGTH: 52
206 <212> TYPE: DNA
207 <213> ORGANISM: Oligonucleotide Primer
209 <400> SEQUENCE: 13
210 catgatgaat tctgatcagg gcgatgtcct cctcgatgtc aggccactcg tc              52
213 <210> SEQ ID NO: 14
214 <211> LENGTH: 97
215 <212> TYPE: DNA
216 <213> ORGANISM: Oligonucleotide Template
218 <400> SEQUENCE: 14
219 gagaagatgg tgctggagaa caacttcttc gtggagacca tgctgcctag caagatcatg      60
221 cgcaagctgg agcctgagga gttcgccgcc tacctgg                                97
224 <210> SEQ ID NO: 15
225 <211> LENGTH: 99
226 <212> TYPE: DNA
227 <213> ORGANISM: Oligonucleotide Template
229 <400> SEQUENCE: 15
230 cttcaccaga gggatctcgc gaggccagct cagggtaggg cggcgcaact cgcccttctc      60
232 cttgaagggc tccaggtagg cggcgaactc ctcaggctc                            99
235 <210> SEQ ID NO: 16

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RAW SEQUENCE LISTING

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DATE: 12/17/2003

TIME: 07:58:39

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12162003\J721405.raw

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236 <211> LENGTH: 62
237 <212> TYPE: DNA
238 <213> ORGANISM: Oligonucleotide Primer
240 <400> SEQUENCE: 16
241 gatacatcta gatgatcaag agcgaggagg gcgagaagat ggtgctggag aacaacttct .60
243 tc 62
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247 <211> LENGTH: 67
248 <212> TYPE: DNA
249 <213> ORGANISM: Oligonucleotide Primer
251 <400> SEQUENCE: 17
252 agttgcgcac gatctgcacc acgtcaggct tgccgccctt caccagaggg atctcgcgag 60
254 gccagct 67
257 <210> SEQ ID NO: 18
258 <211> LENGTH: 100
259 <212> TYPE: DNA
260 <213> ORGANISM: Oligonucleotide Template
262 <400> SEQUENCE: 18
263 cgcgccagcg acgacctgcc caagatgttc atcgagagcg accctggctt cttcagcaac 60
265 gccatcgtgg agggcgccaa gaagttccct aacaccgagt 100
268 <210> SEQ ID NO: 19
269 <211> LENGTH: 100
270 <212> TYPE: DNA
271 <213> ORGANISM: Oligonucleotide Template
273 <400> SEQUENCE: 19
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276 tcaccttcac gaactcgggt ttagggaact tcttggcgcc 100
279 <210> SEQ ID NO: 20
280 <211> LENGTH: 69
281 <212> TYPE: DNA
282 <213> ORGANISM: Oligonucleotide Primer
284 <400> SEQUENCE: 20
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287 cccaagatg 69
290 <210> SEQ ID NO: 21
291 <211> LENGTH: 72
292 <212> TYPE: DNA
293 <213> ORGANISM: Oligonucleotide Primer
295 <400> SEQUENCE: 21
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298 gccatctcg tc 72
301 <210> SEQ ID NO: 22
302 <211> LENGTH: 22
303 <212> TYPE: DNA
304 <213> ORGANISM: Primers used for Mutagenesis
306 <400> SEQUENCE: 22
307 cctctgtatc atatatgctt ta 22
310 <210> SEQ ID NO: 23
311 <211> LENGTH: 22

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RAW SEQUENCE LISTING

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DATE: 12/17/2003

TIME: 07:58:39

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12162003\J721405.raw

312 <212> TYPE: DNA

313 <213> ORGANISM: Primers used for Mutagenesis

315 <400> SEQUENCE: 23

316 taaagcatat atgatacaga gg

22

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/721,405

DATE: 12/17/2003

TIME: 07:58:40

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\12162003\J721405.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number